

Supplementary files

Table S1. Primers sequences, target genes, amplicon sizes and cycling conditions for virulence factors

Gene name	Location/Function	Primers sequences	Amplified segment (bp)	Primary denaturation	Amplification (35 cycles)			Final extension	Reference
					Secondary denaturation	Annealing	Extension		
<i>invA</i>	*SPI-1/Invasion of macrophages	GTGAAATTATGCCACGTTCG GGCAA	284	94°C 5 min.	94°C 30 sec.	55°C 30 sec.	72°C 30 sec.	72°C 7 min.	Oliveira et al. [6]
<i>csgD</i>	Chromosome/master regulator of the biofilm matrix compounds	TTACCGCCTGAGATTATCGT ATGTTAACGAGTCATAG	651	94°C 5 min.	94°C 30 sec.	50°C 45 sec.	72°C 45 sec.	72°C 10 min.	Bhowmick et al. [41]
<i>MgtC</i>	SPI-3/ Mg ²⁺ uptake	TGA CTA TCA ATG CTC CAG TGA AT ATT TAC TGG CCG CTA TGC TGT TG		94°C 5 min.	94°C 45 sec.	58°C 45 sec.	72°C 45 sec.	72°C 10 min.	
<i>bcfC</i>	Chromosome/Bovine colonisation factor, fimbrial usher	ACC AGA GAC ATT GCC TTC C TTC TGC TCG CCG CTA TTC G	467	94°C 5 min.	94°C 45 sec.	53°C 45 sec.	72°C 45 sec.	72°C 10 min.	Huehn et al. [42]
<i>sopE1</i>	Cryptic bacteriophage/Promotes membrane ruffling and disrupts tight junctions	ACT CCT TGCACA ACC AAA TGC GGA TGT CTTCTG CAT TTC GCC ACC		94°C 5 min.	94°C 45 sec.	58°C 45 sec.	72°C 45 sec.	72°C 10 min.	
<i>avrA</i>	SPI-1/Controls <i>Salmonella</i> induced	CCT GTA TTG TTG AGC GTC TGG	422	94°C 5 min.	94°C 45 sec.	58°C 45 sec.	72°C 45 sec.	72°C 10 min.	

	Inflammation	AGA AGA GCT TCG TTG AAT GTC C							
<i>pefA</i>	Plasmid/Plasmid encoded fimbriae	TGT TTC CGG GCT TGT GCT	700	94°C 5 min.	94°C 45 sec.	55°C 45 sec.	72°C 45 sec.	72°C 10 min.	Murugkar et al. [43]
		CAG GGC ATT TGC TGA TTC TTC C							
<i>Stn</i>	Chromosome/ Enterotoxin	TTG TGT CGC TAT CAC TGG CAA CC	617	94°C 5 min.	94°C 45 sec.	59°C 45 sec.	72°C 45 sec.	72°C 10 min.	
		ATT CGT AAC CCG CTC TCG TCC							
<i>hilC</i>	SPI-1/ modulate invasion gene expression	GGACTTGTGCCAGGGATG	241	94°C 5 min.	94°C 30 sec.	62°C 30 sec.	72°C 30 sec.	72°C 7 min.	Yang et al. [44]
		TGACCATTGCGGGTGAG							
<i>ompF</i>	Chromosome/ Outer membrane porins	CCTGGCAGCGGTGATCC	519	94°C 5 min.	94°C 45 sec.	50°C 45 sec.	72°C 45 sec.	72°C 10 min.	Tatavart hy and Cannons [23]
		TGGTGTAACCTACGCCATC							

*SPI: *Salmonella* Pathogenicity Island

Table S2. Distribution of virulence genes combinations and antibiotic resistance patterns in the different *Salmonella* serovars

Serovars (n)	Sample	Antibiotic resistance (pattern profile)	Genetic profile	Virulence genes
<i>S. Enteritidis</i> (2)	Cloacal swabs	SXT,AMP,AMC,CEC,FEP,CTX	P2	<i>invA, csgD, bcfC, stn, avrA, mgtC, ompF</i>
<i>S. Enteritidis</i> (2)	Cloacal swabs	SXT, AMP,S,DO,CIP	P2	<i>invA, csgD, bcfC, stn, avrA, mgtC, ompF</i>
<i>S. Enteritidis</i> (2)	Cloacal swabs	SXT,AMC	P4	<i>invA, csgD, hilC, bcfC, stn</i>
<i>S. Enteritidis</i> (2)	Cloacal swabs	SXT, AMP,AMC,S, DO,CEC	P4	<i>invA, csgD, hilC, bcfC, stn</i>
<i>S. Enteritidis</i> (2)	Cloacal swabs	SXT, AMP,AMC,S,DO	P5	<i>invA, csgD, hilC</i>
<i>S. Enteritidis</i> (4)	Cloacal swabs	SXT, AMP,S,DO,CIP	P5	<i>invA, csgD, hilC</i>
<i>S. Enteritidis</i> (6)	Cloacal swabs	SXT, AMP,AMC,S,DO	P3	<i>invA, csgD, hilC, bcfC, stn, avrA, mgtC</i>
<i>S. Enteritidis</i> (6)	Water	SXT, AMP,AMC,S, DO,CEC,C	P1	<i>invA, csgD, hilC, bcfC, stn, avrA, mgtC, ompF</i>
<i>S. Enteritidis</i> (4)	Feed	SXT,AMC	P6	<i>invA, hilC</i>
<i>S. Enteritidis</i> (2)	Carcass	SXT,AMP,AMC,S,CEC,C,FEP	P1	<i>invA, csgD, hilC, bcfC, stn, avrA, mgtC,ompF</i>
<i>S. Enteritidis</i> (2)	Carcass	SXT, AMP,AMC,S,DO	P1	<i>invA, csgD, hilC, bcfC, stn, avrA, mgtC,ompF</i>
<i>S. Enteritidis</i> (2)	Carcass	SXT, AMP,AMC,S, DO,CEC,C	P3	<i>invA, csgD, hilC, bcfC, stn, avrA, mgtC</i>
<i>S. Enteritidis</i> (4)	Carcass	SXT,AMP,AMC,CEC,FEP,CTX	P3	<i>invA, csgD, hilC,bcfC,stn, avrA, mgtC</i>

<i>S. Enteritidis</i> (8)	Carcass	SXT,AMC	P7	<i>invA, csgD</i>
<i>S.Typhimurium</i> (10)	Cloacal swabs	SXT,S, DO,CEC	P6	<i>invA, hilC</i>
<i>S.Typhimurium</i> (9)	Cloacal swabs	SXT,AMC	P7	<i>invA, csgD</i>
<i>S.Typhimurium</i> (4)	Feed	SXT,AMP,AMC,C	P6	<i>invA, hilC</i>
<i>S.Typhimurium</i> (2)	Letter	SXT,AMP,AMC,S,DO,CTX,CI P	P2	<i>invA, csgD, bcfC, stn, avrA, mgtC, ompF</i>
<i>S.Typhimurium</i> (3)	Carcass	SXT,S, DO,CEC	P2	<i>invA, csgD, bcfC, stn, avrA, mgtC, ompF</i>
<i>S. Typhimurium</i> (2)	Carcass	SXT,AMP,AMC,S,CEC,C,FEP	P3	<i>invA, csgD, hilC, bcfC, stn, avrA, mgtC</i>
<i>S.Typhimurium</i> (2)	Carcass	SXT, AMP,AMC,S, DO,CEC	P4	<i>invA, csgD, hilC, bcfC, stn</i>
<i>S.Typhimurium</i> (1)	Carcass	SXT,AMP,S	P4	<i>invA, csgD, hilC, bcfC, stn</i>
<i>S.Typhimurium</i> (2)	Carcass	SXT,AMP,AMC,CEC,FEP,CTX	P5	<i>invA, csgD, hilC</i>
<i>S.Typhimurium</i>	Carcass	SXT,S, DO,CEC	P6	<i>invA, hilC</i>
<i>S.Kentucky</i>	Cloacal swabs	SXT, AMP,AMC,S,DO	P7	<i>invA, csgD</i>
<i>S.Kentucky</i> (2)	Carcass	SXT, AMP,AMC,S, DO,CEC,C	P5	<i>invA, csgD, hilC</i>
<i>S.Kentucky</i> (2)	Carcass	SXT,AMP,S	P7	<i>invA, csgD</i>
<i>S.Kentucky</i> (2)	Carcass	SXT,AMP,AMC,CEC,FEP,CTX	P7	<i>invA, csgD</i>
<i>S.Kentucky</i>	Carcass	SXT,AMP,AMC,S,DO,CTX,CI P	P5	<i>invA, csgD, hilC</i>

<i>S.Kentucky</i>	Carcass	SXT,AMP,AMC,S,DO,CTX,CI P	P5	<i>invA, csgD, hilC</i>
<i>S.Kentucky</i> (3)	Fan swabs	SXT, AMP,AMC,S,DO	P7	<i>invA, csgD</i>
<i>S.Molade</i> (6)	Cloacal swabs	SXT,AMP,S	P5	<i>invA, csgD, hilC</i>
<i>S.Molade</i> (2)	Carcass	SXT, AMP,AMC,S, DO,CEC	P7	<i>invA, csgD</i>
<i>S.Molade</i> (4)	Carcass	SXT,AMC	P7	<i>invA, csgD</i>
<i>S. Bargny</i> (2)	Carcass	SXT,AMP,S	P5	<i>invA, csgD, hilC</i>
<i>S. Bargny</i> (2)	Cloacal swab	SXT,AMP,S	P5	<i>invA,csgD, hilC</i>
<i>S. Bargny</i> (2)	Water	SXT,AMC	P7	<i>invA, csgD</i>
<i>S. Inganda</i> (4)	Carcass	SXT,AMP,S	P7	<i>invA, csgD</i>
<i>S. Infantis</i>	Carcass	SXT,AMP,S	P7	<i>invA, csgD</i>
<i>S. Infantis</i>	Workers hand swabs	SXT,AMP,S	P7	<i>invA, csgD</i>

AMC=Amoxicillin-clavulanic acid, AMP=ampicillin, CEC=Cefaclor, CTX=Cefotaxime, FEP=Cefepime, DO=Doxycycline, CIP=Ciprofloxacin, IPM=Imipenem, S=Streptomycin, C=Chloramphenicol, and SXT= trimethoprim sulfamethoxazole.

